Appl. No. 10/049,569 Amdt. dated November 29, 2004 Reply to Office Action of, May 27, 2004

REMARKS

Support for Claim 12 can be found in the specification, e.g., Page 9, lines 31-37.

- 3) Applicants will amend the specification in due course when allowable subject matter is indicated.
- 4)-5). The objection to Claims 6 and 7 was not understood these claims are are directed to different classes of subject matter, and therefore do not appear to be duplicative of other claims. These claims were amended for reasons of clarity, however, and such amendment do not change the scope of the claim.

Claim 8 was amended in the Preliminary amendment filed February 15, 2002 to eliminate the multiple dependency. Therefore, it should not have been withdrawn from consideration. It is currently amended to clarify it. These amendments do not change its scope in any way.

6)-8). The polynucleotides and polypeptides of the present invention can be used as reagents for the detection of gene expression. Such expression has been identified in various organs and tissues, including human brain, kidney, blood, lung, colon, lymph nodes, liver and placenta. See, e.g., Specification, Page 15, lines 29-30. Detection can be performed directly on these tissues, as well as on other specimens, including, urine, biopsy tissue, or autopsy material. See, e.g., Specification, Page 12, lines 23-31. Antibodies to the claimed proteins can be used for a variety of purposes, including for detecting expression in normal and disease states, and as a general marker, e.g., in toxicology experiments to determine whether expression is perturbed by a potentially toxic agent.

Most recent published work by Kilk et al. (Neuropeptides. 2004 Oct; 38(5):316-24), indicates that human galanin receptor type 1 (GalR1) mRNA has been used to optimize antisense

Appl. No. 10/049,569 Amdt. dated November 29, 2004 Reply to Office Action of, May 27, 2004

efficacy and Kofler et al. demonstrate the expression of neuropeptide galanin and galanin receptors in human skin (J Invest Dermatol. 2004 April; 122(4): 1050-3). Page 8, lines 12-15 of the present specification indicates that PGPCR-3 protein is homologous to human galanin receptor.

PGPCR-3 was subsequently re-named GPR-78, and has a separate entry in the On-line Mendelian inheritance in Man (OMIM). Thus, the scientific community generally recognizes it as a useful gene. See Exhibit A. (The attached BLAST search shows 100% identity to SEQ ID NO:2 and GPR-78).

Thus, the Examiner's objections with respect to a lack of utility are not justified.

9). The claims have amended to address the alleged indefiniteness. These amendments do not change the scope of the claims in any way, but merely clarify them.

In view of the above remarks, favorable reconsideration is courteously requested. If there are any remaining issues which could be expedited by a telephone conference, the Examiner is courteously invited to telephone counsel at the number indicated below.

Appl. No. 10/049,569 Amdt. dated November 29, 2004 Reply to Office Action of, May 27, 2004

The Commissioner is hereby authorized to charge any fees associated with this response or credit any overpayment to Deposit Account No. 13-3402.

Respectfully submitted,

Richard M. Lebovitz, Reg. No. 37,067 Attorney for Applicant(s)

MILLEN, WHITE, ZELANO & BRANIGAN, P.C. Arlington Courthouse Plaza 1, Suite 1400 2200 Clarendon Boulevard Arlington, Virginia 22201 Telephone: (703) 243-6333

Facsimile: (703) 243-6410

Attorney Docket No.: MERCK-2378

Date: November 29, 2004

OMIM Online Mendelian Inheritance in Man Johns Hopkins University								
PubMed	1	Nucleotide	Protein	Genome	Structure	PMC	Taxonomy	ОМІМО
Search	ОМІМ		[▼] for				Clear	
Lim	iits	Preview/I	ndex	History	Clipbo	ard I	Details	
Display	Detai	led	Sho	w: 20 💌	Send to To	ext		

*606921 G PROTEIN-COUPLED RECEPTOR 78; GPR78

Gene map locus Chr.4

TEXT

DESCRIPTION

G protein-coupled receptors (GPCRs, or GPRs) contain 7 transmembrane domains and transduce extracellular signals through heterotrimeric G proteins.

CLONING

Lee et al. (2001) identified GPR78 in a genomic database using the sequence of GPR26 (604847) as query. PCR primers were designed to amplify GPR78 from a genomic library, and overlapping fragments of partial sequences were joined to obtain the full-length cDNA. GPR78 encodes a deduced 363-amino acid protein that shares 56% sequence identity with GPR26 in the transmembrane region. Northern blot analysis revealed a 1.1-kb transcript in pituitary, and 1.1- and 4.2-kb transcripts in placenta. No expression was detected in brain, skeletal muscle, lung, heart, liver, pancreas, or kidney.

GENE FUNCTION

Van Laar et al. (2000) determined that GPR78 expression increased in fibroblasts or HeLa cells following UV-A irradiation, exposure to DNA-alkylating agents, or endoplasmic reticulum (ER) stress caused by osmotic shock or the glycosylation inhibitor tunicamycin. The response depended upon the cell line studied. UV-B was a weaker inducer, and UV-C and several other DNA-damaging agents did not induce GPR78 expression. Induction of GPR78 by tunicamycin required activation of multiple ER stress-response elements in the promoter of the GPR78 gene, and induction by a DNA-alkylating agent was independent of the unfolded protein response.

MAPPING

Lee et al. (2001) mapped the GPR78 gene to chromosome 4 based on sequence similarity between the GPR78 sequence and a genomic clone (GenBank AC007104) localized to chromosome 4.

REFERENCES

11/29/2004 12:34 PN

Links

1. Lee, D. K.; Nguyen, T.; Lynch, K. R.; Cheng, R.; Vanti, W. B.; Arkhitko, O.; Lewis, T.; Evans, J. F.; George, S. R.; O'Dowd, B. F.:

Discovery and mapping of ten novel G protein-coupled receptor genes. *Gene* 275: 83-91, 2001.

PubMed ID: 11574155

2. van Laar, T.; Schouten, T.; Hoogervorst, E.; van Eck, M.; van der Eb, A. J.; Terleth, C.: The novel MMS-inducible gene Mif1/KIAA0025 is a target of the unfolded protein response pathway. FEBS Lett. 469: 123-131, 2000.

PubMed ID: 10708769

CONTRIBUTORS

Patricia A. Hartz - updated: 9/2/2003

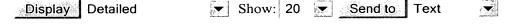
CREATION DATE

Patricia A. Hartz: 5/9/2002

EDIT HISTORY

mgross: 9/2/2003 carol: 5/9/2002

Copyright © 1966-2004 Johns Hopkins University



<u>Disclaimer</u> | <u>Write to the Help Desk</u> | <u>Privacy Policy</u> NCBI | NLM | NIH

```
AN OFFICE TATACCTCGATCGATC
                      2007045641
                      DECAD 40 🕈
                                                                                             Books
                                                Genome
                                                            Structure
                                                                                Taxonomy
                        Nucleotide
                                     Protein
                                                                                  Clear
 Search Nucleotide
                            ▼ for
                                                                             Go
                                                                                         Details
                    Limits
                                      Preview/Index
                                                         History
                                                                        Clipboard
                                                                                         Features
 Display
         default

▼ Show: 20 ▼
                                             Send to
                                                                •
                                                                       Get Subsequence
1: NM 080819. Reports Homo sapiens G pr...[gi:36951033]
                                                                                               Links
                                                                       PRI 23-AUG-2004
             NM 080819
                                       1955 bp
                                                   mRNA
                                                            linear
LOCUS
             Homo sapiens G protein-coupled receptor 78 (GPR78), mRNA.
DEFINITION
             NM 080819
ACCESSION
VERSION
             NM 080819.2 GI:36951033
KEYWORDS
SOURCE
             Homo sapiens (human)
  ORGANISM
             Homo sapiens
             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
             Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE
                (bases 1 to 1955)
             Clark, H.F., Gurney, A.L., Abaya, E., Baker, K., Baldwin, D., Brush, J.,
  AUTHORS
             Chen, J., Chow, B., Chui, C., Crowley, C., Currell, B., Deuel, B.,
             Dowd, P., Eaton, D., Foster, J., Grimaldi, C., Gu, Q., Hass, P.E.
             Heldens, S., Huang, A., Kim, H.S., Klimowski, L., Jin, Y., Johnson, S.,
             Lee, J., Lewis, L., Liao, D., Mark, M., Robbie, E., Sanchez, C.,
             Schoenfeld, J., Seshagiri, S., Simmons, L., Singh, J., Smith, V.
             Stinson, J., Vagts, A., Vandlen, R., Watanabe, C., Wieand, D., Woods, K., Xie, M.H., Yansura, D., Yi, S., Yu, G., Yuan, J., Zhang, M., Zhang, Z.,
             Goddard, A., Wood, W.I., Godowski, P. and Gray, A.
  TITLE
             The secreted protein discovery initiative (SPDI), a large-scale
             effort to identify novel human secreted and transmembrane proteins:
             a bioinformatics assessment
             Genome Res. 13 (10), 2265-2270 (2003)
  JOURNAL
   PUBMED
             12975309
REFERENCE
                (bases 1 to 1955)
  AUTHORS
             Lee, D.K., Nguyen, T., Lynch, K.R., Cheng, R., Vanti, W.B., Arkhitko, O.,
             Lewis, T., Evans, J.F., George, S.R. and O'Dowd, B.F.
             Discovery and mapping of ten novel G protein-coupled receptor genes
  TITLE
  JOURNAL
             Gene 275 (1), 83-91 (2001)
             11574155
   PUBMED
COMMENT
             VALIDATED REFSEQ: This record has undergone preliminary review of
             the sequence, but has not yet been subject to final review. The
             reference sequence was derived from \underline{AK128807.1} and \underline{BC057778.1}.
             On Sep 29, 2003 this sequence version replaced gi: 18201873.
             Summary: G protein-coupled receptors (GPCRs, or GPRs) contain 7
             transmembrane domains and transduce extracellular signals through
             heterotrimeric {\tt G} proteins.[supplied by {\tt OMIM}].
FEATURES
                       Location/Qualifiers
                       1..1955
     source
                       /organism="Homo sapiens"
                       /mol type="mRNA"
                       /db_xref="taxon:9606"
                       /chromosome="4"
                       /map="4p16.1"
     gene
                       1..1955
                       /gene="GPR78"
                       /db xref="GeneID:27201"
                       /db xref="LocusID:27201"
                       /db_xref="MIM:606921"
     CDS
                       420..1511
                       /gene="GPR78"
                       /note="go component: integral to membrane [goid 0016021]
                       [evidence IEA];
                       go_function: rhodopsin-like receptor activity [goid
                       0001584] [evidence IEA];
                       go process: G-protein coupled receptor protein signaling
                       pathway [goid 0007186] [evidence IEA] "
                       /codon start=1
```

```
/product="G protein-coupled receptor 78"
/protein_id="NP 543009.2"
/db_xref="GI:36951034"
/db_xref="GeneID:27201"
/db_xref="LocusID:27201"
/db_xref="LocusID:27201"
/db_xref="MIM:606921"
/translation="MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDMPFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRPRYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRPPEPERPRFAAFTATLHAVGFVLPLAVLCLTSLQVHRVARRHCQRMDTVTMKALALLADLHPSVRQRCLIQQKRRRHRATRKIGIAIATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRPFRQVLAGMVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQQTH"
```

ORIGIN

```
1 acagaagcgc gcagagtccc atcctgccac gccacgagga gagaagaagg aaagatacag
       61 tgttaggaaa gagaceteee tegeeectae geeeegegee eetgegeete getteageet
      121 caggacagte etgeegggae ggtgagegea tteageacee tggacageae egeggttgeg
      181 ctgcctccag ggcggccccg ggctgctcct gctccgcaga gctacgccct cccccgggt
      241 geologgace etgeacttge egelgettte etegegetge tetggacett getagelegge
      301 tetgeacete ecagaageeg tgggegegee geteagetge tecategeet eaettteeea
      361 ggctcgcgcc cgaagcagag ccatgagaac cccagggtgc ctggcgagcc gctagcgcca
      421 tgggccccgg cgaggcgctg ctggcgggtc tcctggtgat ggtactggcc gtggcgctgc
      481 tatccaacgc actggtgctg ctttgttgcg cctacagcgc tgagctccgc actcgagcct
      541 caggogteet eetggtgaat etgtetetgg gecacetget getggeggeg etggacatge 601 cetteacget geteggtgtg atgegegge ggacacegte ggegeeegge geatgecaag
      661 tcattggctt cctggacacc ttcctggcgt ccaacgcggc gctgagcgtg gcggctga
      721 gegeagacea gtggetggea gtgggettee caetgegeta egeeggaege etgegaeege
      781 gctatgccgg cctgctgctg ggctgtgcct ggggacagtc gctggccttc tcaggcgctg
841 cacttggctg ctcgtggctt ggctacagca gcgccttcgc gtcctgttcg ctgcgcctgc
      901 egecegagee tgagegteeg egettegeag cetteacege caegetecat geegtggget
      961 tegtgetgee getggeggtg etetgeetea eetegeteea ggtgeacegg gtggeacgea
     1021 gacactgcca gcgcatggac accgtcacca tgaaggcgct cgcgctgctc gccgacctgc
     1081 accocagtgt geggeagege tgeeteatee ageagaageg gegeegeeae egegeeaeea
     1141 ggaagattgg cattgctatt gcgaccttcc tcatctgctt tgccccgtat gtcatgacca
     1201 ggctggcgga gctcgtgccc ttcgtcaccg tgaacgccca gtggggcatc ctcagcaagt
     1261 gcctgaccta cagcaaggcg gtggccgacc cgttcacgta ctctctgctc cgccggccgt
1321 tccgccaagt cctggccggc atggtgcacc ggctgctgaa gagaaccccg cgcccagcat
     1381 ccacccatga cagctetetg gatgtggccg gcatggtgca ccagctgctg aagagaaccc
     1441 egegeceage gtecacecae aacggetetg tggacacaga gaatgattee tgeetgeage
     1501 agacacactg agggcctggc agggctcatc gccccacct tctaagaagc cctgtggaaa
     1561 gggcactggc cctgccacag agatgccact ggggaccccc agacaccagt ggcttgactt
     1621 tgagctaagg ctgaagtaca ggaggaggag gaggagggg ccggatgtgg gtgtggacag
     1681 cagtagtggc ggaggagagc tcggggctgg gctgcctggc tgctgggtgg ccccgggaca
     1741 gtggcttttc ctctctgaac cttagcttcc tcacccttgt tctggggtca tggcgatgct
     1801 tegagacagt gggtagggaa gtgccetgtg tggcatatgg tactegtggg cgtgctataa
     1861 gtgactgctg ttcatgtggg tgaggtggtc actcttgctc agggtctgtt gtgcagccca
     1921 gatggacacc tgtttctcca aaaaaaaaa aaaaa
//
```

Disclaimer | Write to the Help Desk NCBI | NLM | NIH

Nov 23 2004 06:38:17

```
⊚⊚⊚
                                                                                              Books
                                                             Structure
                                      Protein
                                                 Genome
                        Nucleotide
 Search Protein
                            ▼ for
                                                                                   Clear
                                                                              Go
                                                                        Clipboard
                                                                                          Details
                    Limits
                                       Preview/Index
                                                         History
                                                                                         Features
 Display GenPept
                        -
                            Show: 20
                                       -
                                             Send to
                                                      File
                                                                       Get Subsequence
                                                                                       BLink, Domains,
1: NP 543009. Reports G protein-coupled...[gi:36951034]
                                                                                                Links
LOCUS
             NP_543009
                                         363 aa
                                                             linear
                                                                       PRI 23-AUG-2004
             G protein-coupled receptor 78 [Homo sapiens].
DEFINITION
ACCESSION
             NP 543009
             NP 543009.2 GI:36951034
VERSION
DBSOURCE
             REFSEQ: accession NM 080819.2
KEYWORDS
SOURCE
             Homo sapiens (human)
             Homo sapiens
  ORGANISM
             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
             Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE
                 (residues 1 to 363)
  AUTHORS
             Clark, H.F., Gurney, A.L., Abaya, E., Baker, K., Baldwin, D., Brush, J.,
             Chen, J., Chow, B., Chui, C., Crowley, C., Currell, B., Deuel, B., Dowd, P., Eaton, D., Foster, J., Grimaldi, C., Gu, Q., Hass, P.E.,
             Heldens, S., Huang, A., Kim, H.S., Klimowski, L., Jin, Y., Johnson, S.,
             Lee, J., Lewis, L., Liao, D., Mark, M., Robbie, E., Sanchez, C.,
             Schoenfeld, J., Seshagiri, S., Simmons, L., Singh, J., Smith, V.
             Stinson, J., Vagts, A., Vandlen, R., Watanabe, C., Wieand, D., Woods, K., Xie, M.H., Yansura, D., Yi, S., Yu, G., Yuan, J., Zhang, M., Zhang, Z.,
             Goddard, A., Wood, W.I., Godowski, P. and Gray, A.
  TITLE
             The secreted protein discovery initiative (SPDI), a large-scale
             effort to identify novel human secreted and transmembrane proteins:
             a bioinformatics assessment
  JOURNAL
             Genome Res. 13 (10), 2265-2270 (2003)
   PUBMED
             12975309
REFERENCE
                 (residues 1 to 363)
             Lee, D.K., Nguyen, T., Lynch, K.R., Cheng, R., Vanti, W.B., Arkhitko, O.,
  AUTHORS
             Lewis, T., Evans, J.F., George, S.R. and O'Dowd, B.F.
             Discovery and mapping of ten novel G protein-coupled receptor genes
  TITLE
  JOURNAL
             Gene 275 (1), 83-91 (2001)
             11574155
   PUBMED
COMMENT
             VALIDATED REFSEQ: This record has undergone preliminary review of
             the sequence, but has not yet been subject to final review. The
             reference sequence was derived from AK128807.1 and BC057778.1.
             On Sep 29, 2003 this sequence version replaced gi:18201874.
             Summary: G protein-coupled receptors (GPCRs, or GPRs) contain 7
             transmembrane domains and transduce extracellular signals through
             heterotrimeric G proteins. [supplied by OMIM].
FEATURES
                       Location/Qualifiers
                       1..363
     source
                       /organism="Homo sapiens"
                       /db xref="taxon:9606"
                        /chromosome="4"
                       /map="4p16.1"
                       1..363
     Protein
                       /product="G protein-coupled receptor 78"
     Region
                       60..294
                       /region name="7 transmembrane receptor (rhodopsin family)"
                       /note="7tm_1"
                       /db xref="CDD:5814"
     CDS
                       1..363
                       /gene="GPR78"
                       /coded by="NM 080819.2:420..1511"
                       /note="go component: integral to membrane [goid 0016021]
                        [evidence IEA];
                       go_function: rhodopsin-like receptor activity [goid
```

<u>Disclaimer | Write to the Help Desk</u> <u>NCBI | NLM | NIH</u>

Nov 23 2004 06:38:17



results of BLAST

BLASTP 2.2.10 [Oct-19-2004]

Reference:

Altschul, Stephen F., Thomas L. Madden, Alejandro A. Schäffer, Jinghui Zhang, Zheng Zhang, Webb Miller, and David J. Lipman (1997), "Gapped BLAST and PSI-BLAST: a new generation of protein database search programs", Nucleic Acids Res. 25:3389-3402.

RID: 1101749500-20677-178263183679.BLASTQ4

Query=

(363 letters)

If you have any problems or questions with the results of this search please refer to the ${\tt BLAST}$ ${\tt FAQs}$

Taxonomy reports

Distribution of 514 Blast Hits on the Query Sequence

| Color Key for filignment Scores | C40 | 40-50 | 50±80 | 30±200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-200 | 2-2

Sequences producing significant alignments:	Score (bits)	E Value
gi 37183331 gb AAQ89465.1 GPR78 [Homo sapiens] >gi 3695103	720	0.0 G
gi 16566319 gb AAL26479.1 G protein-coupled receptor [Homo	714	0.0
gi 55622242 ref XP 526521.1 PREDICTED: similar to G protei	435	e-120
gi 50747354 ref XP 426354.1 PREDICTED: similar to G protei	347	3e-94 G
gi 23592220 ref NP 703143.1 G protein-coupled receptor 26	330	5e-89 🕒
gi 20301978 ref NP_620196.1 G protein-coupled receptor 26	321	2e-86 G
gi 27734158 ref NP 775586.1 G protein-coupled receptor 26	320	3e-86 G
gi 50749947 ref XP 421809.1 PREDICTED: similar to G protei gi 47222325 emb CAG05074.1 unnamed protein product [Tetrao	$\frac{315}{301}$	9e-85 🛂 2e-80
gi 47220248 emb CAG03282.1 unnamed protein product [Tetrao	295	1e-78
gi 55634715 ref XP 521629.1 PREDICTED: similar to G protei	<u>241</u>	2e-62
gi 26335307 dbj BAC31354.1 unnamed protein product [Mus mu	<u>197</u>	3e-49 G
gi 29611572 gb AAO85097.1 G protein-coupled receptor GPR26 gi 47214058 emb CAG00716.1 unnamed protein product [Tetrao	$\frac{189}{188}$	le-46 G 2e-46
gi 47215666 emb CAG04750.1 unnamed protein product [Tetrao	95	2e-18
gi 2143560 pir I57942 5-hydroxytryptamine receptor - rat >	90	8e-17 G
gi 50738634 ref XP 426102.1 PREDICTED: similar to Somatostgi 47220082 emb CAG12230.1 unnamed protein product [Tetrao	<u>89</u> 89	2e-16
gi 1162962 gb AAA92633.1 5-HT6 serotonin receptor	89	2e-16
gi 20302609 gb AAM18805.1 type five-like somatostatin rece gi 37499136 gb AAQ91625.1 dopamine D1/beta receptor [Branc	<u>88</u> 87	4e-16 5e-16
gi 13242259 ref NP_077341.1 5-hydroxytryptamine (serotonin	87	7e-16 🕒
gi 38016883 gb AAR07900.1 5-hydroxytryptamine/serotonin re	87	9e-16 G
gi 55586381 ref XP 524584.1 PREDICTED: 5-hydroxytryptamine	87	9e-16
gi 9794865 gb AAF98367.1 somatostatin receptor type two [C	87	9e-16 1e-15
gi 37729012 gb AA003561.1 somatostatin receptor subtype 5	86	
gi 46575715 gb AAH69063.1 Somatostatin receptor 4 [Homo sa gi 55651326 ref XP 525282.1 PREDICTED: similar to dJ753D10	86	2e-15
gi 10946684 ref NP 067333.1 5-hydroxytryptamine (serotonin gi 2340855 emb CAA74971.1 D1A4 Dopamine receptor [Cyprinus	86	2e-15
gi 4557863 ref NP 001043.1 somatostatin receptor 4 [Homo s gi 3941549 gb AAC82382.1 putative odorant receptor LOR4 [L	<u>85</u> 85	3e-15
gi 5689779 emb CAB51953.1 dJ753D10.1 (somatostatin recepto	84	6e-15 G
gi 20899638 ref XP 139909.1 somatostatin receptor 5 [Mus m	84	6e-15 📴
gi 8101115 gb AAF72547.1 somatostatin subtype 5 receptor [84	6e-15 🖪
gi 16945894 gb AAL32173.1 somatostatin receptor 2 [Takifug	84	6e-15
gi 2623672 gb AAB86492.1 somatostatin receptor type 5 [Mus	84	6e-15 📴
gi 1359759 emb CAA66832.1 histamine H2 receptor [Homo sapi	84	8e-15 G
gi 50755091 ref XP 425208.1 PREDICTED: similar to histamin gi 2340857 emb CAA74973.1 DIB Dopamine receptor [Cyprinus	84	8e-15 8e-15
gi 21684966 emb CAD29615.1 somatostatin receptor subtype 5	84	8e-15 G
gi 1085593 pir JC4120 histamine H2 receptor - guinea pig > gi 2119498 pir I51660 dopamine D1B receptor - African claw	<u>83</u> 83	1e-14 1e-14
gi 6680325 ref NP_032339.1 5 hydroxytryptamine receptor 4	83	1e-14 G
gi 3646355 emb CAA09598.1 serotonin 4 receptor [Mus musculus]	83	1e-14 😉
gi 3647303 emb CAA70776.1 serotonin 4 receptor [Mus musculus]	83	1e-14 😘
gi 3647301 emb CAA70775.1 serotonin 4 receptor [Mus musculus]	83	1e-14 😘
gi 47219388 emb CAG01551.1 unnamed protein product [Tetrao	82	2e-14
gi 46451435 gb AAS97962.1 type 2 somatostatin receptor [As	82	2e-14 2e-14
gi 2062423 gb AAC53353.1 somatostatin receptor type 5 [Mus	<u>82</u> 82	3e-14 B
gi 6981588 ref NP 037014.1 somatostatin receptor 5 [Rattus gi 3183689 emb CAA73912.1] serotonin receptor 4 [Cavia porc	82	3e-14 3e-14
gi 1204090 emb CAA56455.1 dopamine receptor [Takifugu rubr	81	4e-14
<u>gi 6978781 ref NP 036900.1 </u> dopamine receptor 5 [Rattus nor	81	4e-14 G

2 of 130

•		
gi 2209143 gb AAB61418.1 somatostatin receptor type 5 [Mus	81	4e-14 😉
gi 3646424 emb CAA09599.1 serotonin 4 receptor [Rattus nor	81	4e-14 G
gi 6446420 gb AAF08613.1 somatostatin receptor type 1 subt	81	4e-14
gi 12274900 emb CAC22248.1 5-hydroxytryptamine4 receptor [81	5e-14 G
gi 29789042 ref NP 038531.1 dopamine receptor 5 [Mus muscu	81	5e-14 G
gi 11321563 ref NP_000861.1 5-hydroxytryptamine (serotonin	81	5e-14 G
gi 41282074 ref NP_955525.1 5-hydroxytryptamine (serotonin	81	5e-14 G
gi 3326989 emb CAA73108.1 5-HT4 receptor [Homo sapiens]	81	5e-14 G
gi 3326991 emb CAA73109.1 5-HT4 receptor [Homo sapiens]	81	5e-14 😉
gi 12274906 emb CAC22251.1 5-hydroxytryptamine4 receptor [81	5e-14 G
gi 40643226 emb CAC79538.1 serotonin receptor 5-HT4 [Homo	81	5e-14 😉
gi 1518034 gb AAC60067.1 dopamine D1A1 receptor gi 3962388 emb CAA06536.1 dopamine D1/beta receptor [Branc	81	5e-14 5e-14
gi 26005719 emb CAD58392.1 5-hydroxytryptamine 4 receptor	81	5e-14 G
gi 50757971 ref XP_425384.1 PREDICTED: similar to Somatost	80	6e-14 G
gi 6900062 emb CAB71316.1 5-HT4 receptor [Homo sapiens]	80	6e-14 G
gi 47222737 emb CAG01704.1 unnamed protein product [Tetrao	80	6e-14
gi 6981584 ref NP 036851.1 somatostatin receptor 1 [Rattus	80	8e-14 🕒
gi 55640843 ref XP 522831.1 PREDICTED: somatostatin recept	80	8e-14 G
gi 54696710 gb AAV38727.1 somatostatin receptor 1 [synthet	80	8e-14 8e-14 G
gi 6678037 ref NP 033242.1 somatostatin receptor 1 [Mus mu gi 6446422 gb AAF08614.1 somatostatin receptor type 1 subt	80	8e-14 5 8e-14
gi 4503391 ref NP_000789.1 dopamine receptor D5 [Homo sapi	80	1e-13 G
gi 2340853 emb CAA74970.1 D1A3 Dopamine receptor (Cyprinus	80	1e-13
gi 89048 pir A39008 histamine H2 receptor - dog >gi 163952	<u>79</u> 79	1e-13 1e-13
gi 29570499 gb AA091738.1 Dopamine receptor protein 1, iso gi 29570498 gb AA091737.1 Dopamine receptor protein 1, iso	79	1e-13
gi 22658483 gb AAN01276.1 dopamine receptor D5 [Homo sapiens]	79	1e-13 G
gi 3941551 gb AAC82383.1 putative odorant receptor LOR14 [79	le-13
gi 47228936 emb CAG09451.1 unnamed protein product [Tetrao	79	1e-13 2e-13 G
gi 14336736 gb AAK61266.1 somatostatin receptor type 5 [Ho	<u>79</u> 79	2e-13 G
gi 4557865 ref NP 001044.1 somatostatin receptor 5 [Homo s gi 2119491 pir 151659 dopamine DlA receptor - African claw	79	2e-13 — 2e-13
gi 2136182 pir 157955 somatostatin receptor - human >gi 43	79	2e-13 😘
gi 55643121 ref XP 510725.1 PREDICTED: somatostatin recept	79	2e-13
gi 49115095 gb AAH72912.1 MGC80373 protein [Xenopus laevis]	<u>79</u> 79	2e-13 5 2e-13
gi 47219262 emb CAG11724.1 unnamed protein product [Tetrao qi 49169818 ref NP 033245.2 somatostatin receptor 4 [Mus m	79	2e-13 2e-13 G
gi 49169818 ref NP 033245.2 somatostatin receptor 4 [Mus m gi 6754260 ref NP 034613.1 5-hydroxytryptamine (serotonin)	79	2e-13 3
gi 6981586 ref NP 037168.1 somatostatin receptor 4 [Rattus	79	2e-13 G
gi 31204165 ref XP_311031.1 ENSANGP00000020010 [Anopheles	79	2e-13 G
gi 31560584 ref NP_034384.2 galanin receptor 2 [Mus muscul	79	2e-13 G
gi 21654945 gb AAL23575.1 putative G-protein coupled recep	79	2e-13
gi 11225272 ref NP_062221.1 somatostatin receptor 2 [Rattu	_78	3e-13 🤁
gi 25990354 gb AAN76495.1 type five somatostatin receptor	_78	3e-13
gi 21314570 gb AAM47010.1 histamine receptor H2 [Mus muscu	<u>78</u>	3e-13 G
gi 12860788 dbj BAB32044.1 unnamed protein product [Mus mu gi 55645903 ref XP_511653.1 PREDICTED: somatostatin recept	<u>78</u> 78	3e-13 G 4e-13
gi 9506709 ref NP 062045.1 galanin receptor 2 [Rattus norv	78	4e-13 G
gi 27806653 ref NP_776467.1 dopamine receptor D1 [Bos taur	78	4e-13 G
gi 50344544 emb CAD59057.1 5-hydroxytryptamine receptor 5A	78	4e-13
gi 11878036 gb AAG40780.1 somatostatin receptor 1 [Sus scr	<u>78</u>	4e-13
gi 4503905 ref NP 003848.1 galanin receptor 2 [Homo sapien	77	5e-13 G 5e-13 G
gi 54696714 gb AAV38729.1 somatostatin receptor 2 [Homo sa gi 456851 gb AAB29143.1 DIA dopamine receptor; DIA recepto	- <u>77</u> - 77	5e-13 5 e-13
gi 543108 pir JC2083 somatostatin receptor 2 - pig >gi 464	77	5e-13
gi 13435405 ref NP 071640.1 histamine receptor H2 [Homo sa	<u>77</u>	5e-13 G
gi 32493367 gb AAH54510.1 HRH2 protein [Homo sapiens]	77	5e-13 G
gi 55625426 ref XP 527128.1 PREDICTED: similar to HRH2 pro	_ 77	5e-13
gi 3642918 gb AAC36589.1 galanin receptor type 2 [Mus musc	77	5e-13 📴

gi 47086925 ref NP_998462.1 zgc:85682 [Danio rerio] >gi 46	<u>77</u>	5e-13 🖪
gi 6680275 ref NP 032312.1 histamine receptor H 2 [Mus mus	77	5e-13 G
gi 50732639 ref XP 425970.1 PREDICTED: similar to 5-hydrox	77	5e-13 G
gi 3941547 gb AAC82381.1 putative odorant receptor LOR3 [L gi 7229404 gb AAF42810.1 somatostatin receptor 2B [Homo sa	$\frac{77}{77}$	5e-13 5e-13
gi 14550544 gb AAH09522.1 Unknown (protein for IMAGE:33547	77	5e-13 G
gi 49902351 gb AAH74796.1 Orexin receptor 1 [Homo sapiens]	77	7e-13 G
gi 1518038 gb AAC60069.1 dopamine D1A2 receptor	77	7e-13
<u>gi 39597901 emb CAE68593.1 </u> Hypothetical protein CBG14463 [gi 2119500 pir I50475 dopamine D1 receptor - goldfish >gi	$\frac{77}{77}$	7e-13 7e-13
gi 178896 gb AAA35550.1 beta-3-adrenergic receptor	77	9e-13 G
gi 4557267 ref NP 000016.1 adrenergic, beta-3-, receptor [77	9e-13 😉
gi 27685697 ref XP 220099.1 similar to putative neurotrans	77	9e-13 😉
gi 55646095 ref XP 523721.1 PREDICTED: similar to galanin	77	9e-13 9e-13
gi 54696712 gb AAV38728.1 somatostatin receptor 2 [synthet gi 4557637 ref NP_001516.1 orexin receptor 1 [Homo sapiens	77	9e-13 G
gi 1070629 pir QRHUB3 beta-3-adrenergic receptor, splice f	77	9e-13
gi 1518036 gb AAC60068.1 dopamine D1C receptor gi 47218969 emb CAG02007.1 unnamed protein product [Tetrao	77	9e-13 9e-13
gi 2144868 pir DYRTD1 dopamine receptor D1 - rat	76	1e-12
gi 34879477 ref XP_341112.1 5-hydroxytryptamine receptor [76	1e-12 G
gi 50754733 ref XP 414481.1 PREDICTED: similar to serotoni	76	1e-12 G
gi 47216965 emb CAG04907.1 unnamed protein product [Tetrao	<u>76</u>	1e-12 1e-12 G
gi 908913 gb AAA70428.1 D1 dopamine receptor protein >gi 1	<u>76</u>	
gi 20857619 ref XP 136992.1 PREDICTED: similar to Putative	76	20 12
gi 6981044 ref NP 037097.1 histamine receptor H 2 [Rattus	76	2e-12 G
gi 13236497 ref NP 076917.1 5-hydroxytryptamine (serotonin qi 47575845 ref NP 001001267.1 serotonin 4A receptor (5-HT	<u>76</u> 76	2e-12 G
gi 47575845 ref NP 001001267.1 serotonin 4A receptor (5-HT gi 27806153 ref NP 776892.1 somatostatin receptor 2 [Bos t	76	2e-12 G
gi 50755087 ref XP 425206.1 PREDICTED: similar to dopamine	76	2e-12 G
gi 6425114 gb AAF08306.1 beta 3 adrenergic receptor [Macac	76	2e-12
gi 45439382 gb AAS18239.2 5-hydroxytryptamine receptor 4 [gi 1362718 pir A55886 dopamine receptor D1A - chicken	76 76	2e-12 G 2e-12
gi 924639 gb AAC52232.1 5-HT4S receptor >gi 1363262 pir S	<u>75</u>	2e-12 3
gi 6981060 ref NP_036985.1 5-hydroxytryptamine (serotonin)	_75	2e-12 G
gi 15558894 emb CAC69545.1 somatostatin receptor subtype 1	75	2e-12 5
gi 4009515 gb AAC95468.1 galanin receptor 2 [Mus musculus]	75 75	3e-12 G
gi 2136496 pir 147217 dopamine receptor - pig >gi 808098 g gi 12643864 sp Q9TT96 B1AR BOVIN Beta-1 adrenergic receptor	75	3e-12 3e-12
gi 4102061 gb AAD01420.1 somatostatin receptor type 2 [Mus	75	3e-12
gi 33859542 ref NP 034206.1 dopamine receptor D1A [Mus mus	<u>75</u>	3e-12 G
gi 48139558 ref XP_397024.1 similar to allatostatin recept	_75	3e-12 G
gi 4102060 gb AAD01419.1 somatostatin receptor type 2 [Mus gi 47219685 emb CAG12607.1 unnamed protein product [Tetrao	75 75	3e-12 15 3e-12
gi 6678039 ref NP 033243.1 somatostatin receptor 2 [Mus mu	74	5e-12 G
gi 55625662 ref XP 527182.1 PREDICTED: dopamine receptor D	74	5e-12 3
gi 55742138 ref NP 001007122.1 5-hydroxytryptamine (seroto	74	5e-12 😘
gi 6680660 ref NP 031442.1 adrenergic receptor, alpha lb [74	5e-12 G
gi 1518040 gb AAC60070.1 dopamine D1B receptor	$\frac{74}{74}$	5e-12
gi 47213581 emb CAF93484.1 unnamed protein product [Tetrao gi 3335678 gb AAC27328.1 D1 dopamine receptor [Macaca mula	74	5e-12 5e-12
gi 30399 emb CAA41734.1 D-1 dopamine receptor [Homo sapiens]	74	5e-12 G
gi 1362720 pir C55886 dopamine receptor D1D - chicken	$\frac{74}{74}$	5e-12 5e-12
gi 49456799 emb CAG46720.1 DRD1 [Homo sapiens] gi 346640 pir S28058 serotonin receptor 5 - mouse	74	6e-12
gi 3283973 gb AAC25414.1 beta 1 adrenergic receptor [Ovis	74	6e-12
gi 50979252 ref NP 001003377.1 beta 3 adrenergic receptor	74	6e-12 G
gi 50755831 ref XP 425241.1 PREDICTED: similar to somatost gi 28827164 gb AAO24755.1 melanin-concentrating hormone re	$\frac{74}{74}$	6e-12 G 6e-12
gi 54638423 gb EAL27825.1 GA21941-PA [Drosophila pseudoobs	74	6e-12
gi 7159252 gb AAF37686.1 octopamine receptor [Aplysia cali	74	6e-12

'gi 47223437 emb CAG04298.1 unnamed protein product [Tetrao	74	6e-12
gi 2137787 pir JC4629 somatostatin receptor type-4 - mouse	74	6e-12 😉
gi 1913918 gb AAB51068.1 beta-3 adrenergic receptor [Canis gi 603869 emb CAA57494.1 D1-like dopamine receptor [Oreoch	$\frac{74}{74}$	6e-12 8e-12
gi 6978775 ref NP 036678.1 dopamine receptor 1A [Rattus no	74	8e-12 G
gi 55241891 gb EAA08140.3 ENSANGP00000018804 [Anopheles ga	74	8e-12
gi 50950129 ref NP 001002933.1 hypocretin receptor 2 [Cani	73	le-11 G
gi 47223876 emb CAG06053.1 unnamed protein product [Tetrao gi 47217696 emb CAG13327.1 unnamed protein product [Tetrao	<u>73</u> 73	le-11 le-11
gi 47214321 emb CAG11192.1 unnamed protein product [Tetrao	73	1e-11
gi 37704009 gb AAR01326.1 orexin receptor type-1 [Mus musc	73	1e-11 G
gi 31746493 gb AAP68899.1 somatostatin receptor type five gi 23379643 gb AAM76564.1 adrenergic receptor beta-3 [Pong	$\frac{73}{73}$	1e-11 1e-11
gi 46451437 gb AAS97963.1 type 3 somatostatin receptor [As	73	1e-11
gi 55665868 emb CAH73407.1 solute carrier family 31 (coppe	72	2e-11
gi 28839657 gb AAH47526.1 HTR7 protein [Homo sapiens] >gi	72	2e-11 G
gi 10880131 ref NP 062874.1 5-hydroxytryptamine receptor 7	72	
gi 10880129 ref NP 062873.1 5-hydroxytryptamine receptor 7 gi 55634399 ref XP 521556.1 PREDICTED: similar to 5-hydrox	72	2e-11
gi 6981018 ref NP 037196.1 hypocretin receptor 1 [Rattus n gi 55664480 emb CAH69965.1 5-hydroxytryptamine (serotonin)	$\frac{72}{72}$	2e-11 2 2e-11
gi 18597350 gb AAL76096.1 somatostatin receptor [Rattus no	<u>72</u> 72	2e-11 G 2e-11
gi 227114 prf 1614340A dopamine receptor D1 gi 2119497 pir 151661 dopamine D1C receptor - African claw	$\frac{72}{72}$	2e-11 2e-11
gi 4557639 ref NP_001517.1 orexin receptor 2 [Homo sapiens	72	2e-11 G
gi 6840859 gb AAF28802.1 octopamine receptor [Aplysia kuro qi 435817 gb AAB28595.1] 5-hydroxytryptamine receptor subty	$\frac{72}{72}$	2e-11 2e-11
gi 435817 gb AAB28595.1 5-hydroxytryptamine receptor subty gi 6680329 ref NP 032341.1 5-hydroxytryptamine (serotonin)	72	3e-11 3
gi 6981062 ref NP_037280.1 5-hydroxytryptamine (serotonin)	72	3e-11 G
gi 55666937 ref XP 528711.1 PREDICTED: similar to D(1B) do	72	3e-11
gi 33329181 gb AAQ09991.1 mu opioid-like receptor [Rana pi	$\frac{72}{72}$	3e-11
gi 3941553 gb AAC82384.1 putative odorant receptor LOR12 [gi 8885888 gb AAF80280.1 alpha 1b adrenoceptor [Oryctolagu	$\frac{72}{72}$	3e-11 3e-11
gi 402163 gb AAA42134.1 5HT-7 serotonin receptor	71	4e-11 G
gi 477007 pir A47519 serotonin receptor 7 - rat >gi 410307	<u>71</u>	4e-11 G
gi 34866769 ref XP_346803.1 hypothetical protein XP_346802	71	4e-11 😘
gi 45387607 ref NP_991152.1 opiate receptor-like [Danio re	71	4e-11 6
gi 47522982 ref NP_999250.1 serotonin 5-hydroxytryptamine	71	4e-11 G
gi 50749576 ref XP_426518.1 PREDICTED: similar to dopamine gi 449413 prf 1919247A betal adrenergic receptor	7 <u>1</u>	4e-11 G 4e-11
gi 50110 emb CAA42966.1 beta-3-adrenergic-receptor [Mus mu	$\frac{71}{71}$	5e-11 G 5e-11
gi 7441613 pir S71323 alpha-1A adrenergic receptor - Japan gi 6678041 ref NP_033244.1 somatostatin receptor 3 [Mus mu	71	5e-11 G
gi 298113 emb CAA51384.1 beta-3-adrenergic receptor [Mus m	71	5e-11 G
gi 55630516 ref [XP_519708.1] PREDICTED: similar to beta-3-a	71	5e-11
gi 32423757 gb AAF97249.2 mu opioid receptor [Macaca mulatta]	71	5e-11
gi 7304871 ref NP 038490.1 adrenergic receptor, beta 3 [Mu	<u>71</u>	5e-11 G
gi 52219136 ref NP 001004654.1 zgc:103757 [Danio rerio] >g	71	50 11 _
gi 50747427 ref XP 420871.1 PREDICTED: similar to alpha 1d	71	50 11 5
gi 1698952 gb AAB37322.1 high-affinity lysophosphatidic ac gi 47213181 emb CAF95370.1 unnamed protein product [Tetrao	71	5e-11 5 5e-11
gi 47207357 emb CAF93600.1 unnamed protein product [Tetrao	71	5e-11
gi 1345417 dbj BAA09921.1 alphalA-adrenoceptor [Oryzias la gi 31746495 gb AAP68900.1 type-three somatostatin receptor	$\frac{71}{70}$	5e-11 7e-11
gi 12621102 ref NP 075227.1 5-hydroxytryptamine (serotonin gi 23379641 gb AAM76563.1 adrenergic receptor beta-3 [Gori	70	7e-11 5 7e-11
<u>gi 51869673 emb CAF31499.1 </u> 5-HT receptor 7a [Canis familia	70	7e-11 G
gi 1857149 gb AAB48396.1 5-hydroxytryptamine7 receptor iso	<u>70</u> 70	7e-11 G 9e-11
gi 2133653 pir S68780 dopamine D1-like receptor - fruit fl gi 627342 pir A55044 beta-4C-adrenergic receptor - turkey	70	9e-11
gi 6978459 ref NP 036833.1 adrenergic receptor, beta 1 [Ra	70	9e-11 😘
gi 34328059 ref NP_038488.1 adrenergic receptor, alpha 1d	70	9e-11 😉

•		_
gi 23171234 gb AAF55030.2 CG9652-PA [Drosophila melanogast	70	9e-11 G
gi 6981020 ref NP_037206.1 hypocretin receptor 2 [Rattus n	70	9e-11 G
gi 6680327 ref NP 032340.1 5-hydroxytryptamine (serotonin)	70	9e-11 G
		9e-11 G
gi 45382489 ref NP 990692.1 Mel-1c melatonin receptor [Gal	70	
gi 6978463 ref NP_037240.1 adrenergic receptor, beta 3 [Ra	70	1e-10 E
gi 220671 dbj BAA00527.1 beta-1 adrenergic receptor [Rattu	_70	le-10 G
gi 2134105 pir I51666 Mel-1c receptor subtype - African cl	70	1e-10
gi 241216 gb AAB20702.1 beta 3-adrenergic receptor [Rattus	70	1e-10 G
gi 6680666 ref NP 031445.1 adrenergic receptor, beta 1 [Mu	70	1e-10 😉
qi 1857131 qb AAB48390.1 Mel-1c(a) melatonin receptor [Xen	70	1e-10 G
gi 32482003 gb AAP84354.1 somatostatin receptor 3 [Homo sa	69	1e-10 G
gi 55586759 ref XP 524646.1 PREDICTED: similar to hypocret	69	1e-10
gi 21951818 gb AAM82355.1 somatostatin receptor type 3 [Ca	69	1e-10
gi 23379645 gb AAM76565.1 adrenergic receptor beta-3 [Sagu	69	1e-10
gi 23379639 gb AAM76562.1 adrenergic receptor beta-3 [Pan	69	1e-10 😘
gi 50747348 ref XP 426351.1 PREDICTED: similar to dopamine	69	1e-10 G
gi 46575616 gb AAH69171.1 Putative neurotransmitter recept	69	1e-10 😉
gi 27695547 gb AAH42068.1 Similar to somatostatin receptor	69	_{le-10} G
	69	2e-10 B
gi 51869675 emb CAF31500.1 5-HT receptor 7b [Canis familia	<u>69</u>	
gi 202764 gb AAA63478.1 alpha-1B adrenergic receptor	69	2e-10 5
gi 54637595 gb EAL26997.1 GA19956-PA [Drosophila pseudoobs gi 6120127 gb AAF04303.1 beta-1 adrenergic receptor [Felis	<u>69</u> 69	2e-10 2e-10
gi 14718772 gb AAK71884.1 mu-opioid receptor [Macaca fasci	69	2e-10
gi 47211074 emb CAF89689.1 unnamed protein product [Tetrao	69	2e-10
gi 1888505 gb AAB53098.1 alpha 1d adrenoceptor [Oryctolagu	69	2e-10
gi 479128 emb CAA54451.1 dopamine receptor [Drosophila mel	69	2e-10
gi 1103944 gb AAA83015.1 5-hydroxytryptamine7 receptor >gi gi 55665067 emb CAH72100.1 RP11-295F4.5 [Homo sapiens]	<u>69</u> 69	2e-10 3e-10
gi 28316758 ref NP_783599.1 G protein-coupled receptor 4 [69	3e-10 G
gi 345733 pir A45121 alpha-1B adrenergic receptor - human	69	3e-10
gi 109444 pir A40491 alpha-1-adrenergic receptor - golden	69	3e-10
gi 26335986 dbj BAC31691.1 unnamed protein product [Mus mu	69	3e-10 G
gi 4501959 ref NP 000670.1 alpha-1B-adrenergic receptor [H	69	3e-10 G
gi 47575853 ref NP 058687.2 adrenergic receptor, alpha 1b	69	3e-10 G
gi 13324696 ref NP_077809.1 adrenergic receptor, alpha 1d	69	3e-10 E
gi 27371132 gb AAH37002.1 Adralb protein [Mus musculus] gi 34368416 emb CAE46112.1 alpha-1B adrenergic receptor [S	<u>69</u> 69	3e-10 5
gi 47213375 emb CAF90994.1 unnamed protein product [Tetrao	69	3e-10
gi 666891 gb AAB59485.1 alpha-1B adrenergic receptor >gi 1	69	3e-10 🖪
gi 547221 gb AAB31164.1 alpha adrenergic receptor subtype	69	3e-10
gi 543734 sp P15823 A1AB_RAT Alpha-1B adrenergic receptor (69	3e-10 😉
gi 37723880 gb AA003563.1 somatostatin receptor subtype 3	68	3e-10 😘
gi 38112417 gb AAR11294.1 orexin receptor type-2a [Mus mus	68	3e-10 G
		3e-10 G
gi 38112416 gb AAR11293.1 orexin receptor type-2b [Mus mus	68	
gi 27806037 ref NP 776833.1 opioid receptor, mu 1 [Bos tau gi 1857135 gb AAB48392.1 Mel-1c(b) melatonin receptor [Xen	<u>68</u> 68	3e-10 🛂 3e-10
gi 3954976 emb CAA06542.1 dopamine Dlx receptor [Myxine gl	68	3e-10
gi 20139232 sp Q9MYW9 OPRM MACMU Mu-type opioid receptor (M	68	3e-10
gi 7690135 gb AAB31163.2 alpha adrenergic receptor subtype	68	4e-10 🕒
gi 86790 pir JH0447 alpha-1A-adrenergic receptor - human >	68	4e-10 G
gi 86529 pir A25896 beta-adrenergic receptor - turkey >gi	68	4e-10
gi 4501957 ref NP_000669.1 alpha-1D-adrenergic receptor [H	68	4e-10 😉
gi 13027456 ref NP 076482.1 orphan G protein-coupled recep	68	4e-10 😘
gi 27446646 gb AAK74189.1 mu opioid receptor variant MOR-1	68	4e-10 G
gi 4588548 gb AAD26148.1 beta 3 adrenergic receptor; beta3	68	4e-10
gi 2661769 emb CAA73841.1 dopamine receptor, D1 [Apis mell	68	4e-10 📴
gi 11128469 emb CAC15482.1 dJ366F13.1 (opioid receptor mu	67	6e-10 G
		6e-10 5
gi 38156309 gb AAR12887.1 mu opioid receptor variant MOR-1	67	06-10 6

gi 4501961 ref NP_000671.1 alpha-1A-adrenergic receptor is	67	6e-10 😉
gi 50959650 gb AAH74927.1 Opioid receptor, mu 1 [Homo sapi	67	6e-10 G
gi 37362413 gb AAQ91331.1 adrenergic alpha 1A receptor [Ho	67	6e-10 G
gi 37729014 gb AA003562.1 somatostatin receptor subtype 4	67	6e-10 G
gi 7300871 gb AAF56012.1 CG6919-PA [Drosophila melanogaste	67	6e-10 G
gi 15451761 ref NP_150647.1 alpha-1A-adrenergic receptor i	67	6e-10 B
gi 15451759 ref NP_150646.1 alpha-1A-adrenergic receptor i	67	6e-10 G
gi 15451757 ref NP_150645.1 alpha-1A-adrenergic receptor i	<u>67</u>	6e-10 G
gi 34850746 ref NP 919242.1 adrenergic, beta-1-, receptor	<u>67</u>	6e-10 G
gi 48374069 ref NP 001001538.1 mu opioid receptor [Sus scr	67	6e-10 G
gi 50801488 ref XP 428541.1 PREDICTED: similar to beta-4C	<u>67</u>	6e-10 G
gi 50759565 ref XP 425762.1 PREDICTED: similar to alpha-1A	67	6e-10 G
gi 50737619 ref XP 426087.1 PREDICTED: similar to opioid r	<u>67</u>	6e-10 G
gi 27446648 gb AAK74190.1 mu opioid receptor variant MOR-1	<u>67</u>	6e-10 G
gi 27373028 gb AAN87342.1 DRG kappa 1 splice variant KOR 1 gi 47217452 emb CAG10221.1 unnamed protein product [Tetrao	<u>67</u> 67	6e-10 G 6e-10
<pre>gi 607912 gb AAB60354.1 mu opioid receptor variant >gi 213</pre>	67	6e-10 G
gi 666893 gb AAB59486.1 alpha-1C-adrenergic receptor	67	6e-10 G
gi 6114881 emb CAB59347.1 alpha-1D adrenergic receptor [Su gi 547222 gb AAB31165.1 alpha adrenergic receptor subtype	<u>67</u> 67	6e-10 6e-10
gi 1362719 pir B55886 dopamine receptor D1B - chicken	67	6e-10
gi 40362763 gb AAR84650.1 alpha 1A adrenoceptor isoform 6 gi 40362761 gb AAR84649.1 alpha 1A adrenoceptor isoform 5b	<u>67</u> 67	6e-10 6e-10
gi 40362757 gb AAR84647.1 alpha 1A adrenoceptor isoform 3c	67	6e-10 G
gi 40362755 gb AAR84646.1 alpha 1A adrenoceptor isoform 3b	67	6e-10 B
gi 40362753 gb AAR84645.1 alpha 1A adrenoceptor isoform 2c	67	6e-10 G
gi 12858052 dbj BAB31185.1 unnamed protein product [Mus mu	67	6e-10 G
gi 55558 emb CAA35934.1 unnamed protein product [Rattus no	67	7e-10 😘
gi 111409 pir S12591 beta-1-adrenergic receptor - rat	67	7e-10
gi 111359 pir A38731 alpha-1A adrenergic receptor - rat >g	67	7e-10 G
gi 28460708 ref NP 783178.1 trace amine receptor 9 [Rattus	67	7e-10 😉
gi 2143857 pir 156517 mu opioid receptor - rat >gi 403574	67	7e-10 E
gi 4505515 ref NP 000905.1 opioid receptor, mu 1 [Homo sap	_67	7e-10 G
gi 32186858 gb AAP72174.1 somatostatin receptor 1 [Canis f	67	7e-10 G
<u>gi 38016137 ref NP_937822.1 </u> G protein-coupled receptor 103 <u>gi 12231866 gb AAG49292.1 </u> 5-hydroxytryptamine 7 receptor [67 67	7e-10 G 7e-10
gi 47117904 sp Q96P65 QRFR HUMAN Orexigenic neuropeptide QR gi 27261706 gb AAN86027.1 mu-opioid receptor [Cavia porcel	67 67	7e-10 5 7e-10
gi 404116 dbj BAA04109.1 kappa opioid receptor [Rattus nor gi 1083836 pir A55259 kappa opioid receptor - guinea pig >	67 67	le-09
gi 425189 gb AAA41496.1 kappa opioid receptor	67	1e-09 G
gi 1204095 emb CAA56457.1 dopamine receptor [Takifugu rubr	67	1e-09 1e-09 13
gi 34866003 ref XP_346733.1 hypothetical protein XP_346732 gi 6981310 ref NP 037203.1 opioid receptor, mu 1 [Rattus n	<u>67</u> 67	1e-09 5
gi 6981310 ref NP 037203.1 opioid receptor, mu 1 [Rattus n gi 35187403 gb AAQ84306.1 type 7 serotonin receptor [Helis	67	1e-09
gi 12232632 gb AAD22540.2 alpha-1A adrenergic receptor [Ca	67	1e-09
gi 37625043 gb AAQ95734.1 dopamine receptor D4 [Mustela pu gi 47217312 emb CAG12520.1 unnamed protein product [Tetrao	<u>67</u> 67	1e-09 1e-09
gi 1657822 gb AAB93648.1 betal adrenergic receptor [Canis	67	1e-09
gi 2143855 pir 156504 mu opioid receptor - rat >gi 1017732 gi 241214 gb AAB20701.1 alpha 1-adrenergic receptor subtyp	<u>67</u> 66	1e-09 📴 1e-09
gi 885865 gb AAA86878.1 mu opioid recptor	66	1e-09 🖪
gi 28212244 ref NP 783177.1 trace amine receptor 6 [Rattus	66	1e-09 📴
gi 39725940 ref NP_000903.2 opioid receptor, kappa 1 [Homo	66	1e-09 G
gi 37724703 gb AA018365.1 mu opioid receptor variant CII [66	1e-09 🖪
gi 6754940 ref NP 035143.1 opioid receptor, mu 1 [Mus musc	66	1e-09 🖪
gi 4505925 ref NP_003958.1 putative neurotransmitter recep	66	1e-09 G
gi 565069 gb AAB60673.1 mu opioid receptor [Mus musculus]	_66	1e-09 💪

•		6
gi 8778198 gb AAF79213.1 mu opioid receptor variant F [Mus	66	1e-09 G
gi 18026695 gb AAL55583.1 mu opioid receptor variant BII [<u>66</u>	1e-09 G
gi 18026693 gb AAL55582.1 mu opioid receptor variant BI [M	_66	1e-09 G
gi 18026691 gb AAL55581.1 mu opioid receptor variant A [Mu	66	1e-09 G
gi 5853309 gb AAD54415.1 mu opioid receptor variant C; MOR	66	1e-09 G
gi 45768619 gb AAH67468.1 G protein-coupled receptor 63 [H	_66	1e-09 G
gi 20379020 gb AAM21070.1 opioid receptor kappa [Homo sapi	<u>66</u>	1e-09 G
gi 27448127 gb AA013794.1 mu opioid receptor variant R [Mu	66	1e-09 G
gi 27448125 gb AA013793.1 mu opioid receptor variant Q [Mu	_66	1e-09 📴
gi 27448123 gb AA013792.1 mu opioid receptor variant P [Mu	66	1e-09 G
gi 27446644 gb AAK74188.1 mu opioid receptor variant MOR-1 gi 1256416 gb AAA96315.1 beta3-adrenergic receptor [Cavia	66 66	1e-09
gi 5805153 gb AAD51861.1 mu opioid receptor MOR1D [Mus mus	66	1e-09 G
gi 3650454 gb AAC61296.1 octopamine receptor type 1 [Lymna	66	1e-09
gi 409029 gb AAA93114.1 alpha1C adrenergic receptor	_66	1e-09 G
gi 26332529 dbj BAC29982.1 unnamed protein product [Mus mu	<u>66</u>	1e-09 G
gi 22832515 gb AAF48875.2 CG6857-PA [Drosophila melanogast	_66	2e-09 G
gi 15004694 gb AAK77197.1 adrenergic receptor alpha-la [Ho gi 47225323 emb CAG09823.1 unnamed protein product [Tetrao	<u>66</u> 66	2e-09 G 2e-09
gi 47213874 emb CAF94024.1 unnamed protein product [Tetrao	66	2e-09
gi 47205254 emb CAF95660.1 unnamed protein product [Tetrao	66	2e-09
gi 47178862 emb CAG13901.1 unnamed protein product [Tetrao	_66	2e-09
gi 32165520 gb AAP72127.1 G protein-coupled receptor 135 [65	2e-09 G
gi 395368 emb CAA49352.1 serotonin receptor [Rattus norveg	65	2e-09 G
gi 1002739 gb AAC50504.1 GPR10 gi 55250889 gb AAH85587.1 Zgc:103685 [Danio rerio] >gi 559	<u>65</u>	2e-09 G 2e-09
gi 54111955 gb AAV28689.1 mu opioid receptor [Taricha gran	65	2e-09
gi 52698314 gb AAR36861.1 melanopsin [Felis catus]	65	2e-09
gi 48101556 ref XP 392683.1 similar to CG4322-PA [Apis mel gi 2796173 gb AAB97525.1 beta-1 adrenergic receptor [Sus s	<u>65</u> 65	2e-09 G 2e-09
gi 2796173 gb AAB97525.1 beta-1 adrenergic receptor [Sus s gi 2398857 dbj BAA22217.1 Gq-coupled rhodopsin [Mizuhopect	65	2e-09
gi 24111248 ref NP 035141.1 opioid receptor, kappa 1 [Mus	65	3e-09 😘
gi 30231226 ref NP 840074.1 opsin 4 (melanopsin) [Danio re	65	3e-09 G
gi 51951314 gb AAU15126.1 kappa opioid receptor [Taricha g	65	3e-09
gi 54642851 gb EAL31595.1 GA16412-PA [Drosophila pseudoobs gi 47222483 emb CAG13003.1 unnamed protein product [Tetrao	<u>65</u> 65	3e-09 3e-09
gi 944892 gb AAB60369.1 dopamine D2 receptor > gi 1706283 s	65	3e-09
<pre>gi 478273 pir JC1525</pre> alpha-1B-adrenergic receptor - rat >g	65	4e-09 😉
gi 18859151 ref NP 571782.1 opioid receptor, mu 1 [Danio r	65	4e-09 🕒
gi 27806213 ref NP_776923.1 adrenergic, alpha 1A, receptor	_65	4e-09 G
gi 50729258 ref XP 425480.1 PREDICTED: similar to Somatost	65	4e-09 🕏
gi 47227683 emb CAG09680.1 unnamed protein product [Tetrao gi 6563386 emb CAB62570.1 alpha-1A adrenergic receptor [Su	<u>65</u> 65	4e-09 4e-09
gi 1438750 gb AAB36304.1 beta 1-adrenergic receptor [Ovis	65	4e-09 G
gi 21928413 dbj BAC05800.1 seven transmembrane helix recep	65	4e-09 B
gi 50749927 ref XP 426540.1 PREDICTED: similar to beta-adr	64	5e-09 G
gi 45708982 gb AAH67455.1 G protein-coupled receptor 45 [H	64	5e-09 G
gi 55627426 ref XP_527507.1 PREDICTED: similar to Putative	64	6e-09
gi 20070983 gb AAH26357.1 G protein-coupled receptor 62 [H	64	6e-09 🕒
gi 3242941 gb AAC23861.1 alpha-1A adrenoreceptor [Canis fa	64	6e-09 📴
gi 2198745 gb AAB61334.1 alpha 1a-adrenoceptor [Oryctolagu gi 8843927 gb AAF80169.1 alpha 1a-adrenoceptor isoform 3 [64 64	6e-09 6e-09
gi 8843927 gb AAF80169.1 alpha la-adrenoceptor isoform 3 [gi 8843925 gb AAF80168.1 alpha la-adrenoceptor isoform 2 [64	6e-09
gi 51765584 ref XP_487102.1 PREDICTED: similar to trace am	64	8e-09 B
gi 31083315 ref NP 009158.3 G protein-coupled receptor 45	64	8e-09 G
gi 37524029 gb AAQ92315.1 relaxin-3 receptor-1 [Homo sapie	64	8e-09 G
gi 31542909 ref NP 444337.2 G protein-coupled receptor 45	64	8e-09 G
gi 55624572 ref XP 526961.1 PREDICTED: G-protein coupled r	64	8e-09
gi 55599359 ref XP 515672.1 PREDICTED: similar to G protei	64	8e-09 8e-09 G
gi 33859500 ref NP 033760.1 adenosine A2a receptor [Mus mu	64	oe-09 ≥

•		
gi 34878896 ref NP 543141.2 G protein-coupled receptor 62	64	8e-09 G 8e-09
gi 23452342 gb AAN33001.1 adenosine-like receptor [Asterin	64	
gi 11993046 gb AAG42572.1 G protein-coupled receptor PSP24 gi 47226910 emb CAG05802.1 unnamed protein product [Tetrao	<u>64</u> 64	8e-09 2 8e - 09
gi 2119488 pir 150081 rhodopsin - green anole >gi 468262 g	63	1e-08
gi 13540557 ref NP 110411.1 G protein-coupled receptor 63	63	1e-08 📴
gi 27683113 ref XP 237112.1 similar to G protein-coupled r	63	1e-08 G
gi 6753710 ref NP 034228.1 opsin (encephalopsin) [Mus musc	63	1e-08 🕒
gi 37497118 ref NP_922917.1 dopamine receptor D2 like [Dan	63	1e-08 📴
gi 41386782 ref NP_776656.1 adrenergic, beta-2, receptor,	63	1e-08 🕒
gi 50742751 ref XP_419740.1 PREDICTED: similar to Vascular	<u>63</u>	1e-08 🕒
gi 45768489 gb AAH67466.1 G protein-coupled receptor 63 [H	63	1e-08
gi 7271779 gb AAF44619.1 rod-like opsin [Salmo salar] gi 6017883 gb AAF01674.1 beta 1 adrenergic receptor [Bos t	<u>63</u> 63	1e-08 1e-08
gi 24432089 ref NP 006047.2 neuromedin U receptor 1 [Homo	63	_{le-08} G
gi 4028154 gb AAC96118.1 putative neurotransmitter recepto	63	1e-08
gi 51765586 ref XP_487103.1 PREDICTED: similar to trace am	62	2e-08 G
gi 2865470 gb AAC02680.1 orphan G protein-coupled receptor	62	2e-08 🕒
gi 30354034 gb AAH51914.1 NMUR1 protein [Homo sapiens] >gi	<u>62</u>	2e-08
gi 33504559 ref NP_878306.1 opioid receptor, kappa 1 [Dani	_62	2e-08 G
gi 45767693 gb AAH67467.1 G protein-coupled receptor 63 [H	<u>62</u> 62	2e-08
gi 22091559 emb CAD23111.1 blue cone opsin [Cottus gobio]	62	2e-08 2e-08 G
gi 4455063 gb AAD21056.1 orphan G protein-coupled receptor	62	2e-08 5
gi 2735351 gb AAB93884.1 high-affinity lysophosphatidic ac gi 71928 pir 000CG rhodopsin - giant octopus	62	2e-08
gi 2695874 emb CAB08107.1 P2Y-like G-protein coupled recep	62	2e-08 😘
gi 808876 dbj BAA06508.1 kappa-opioid receptor [Mus muscul	62	2e-08 G
gi 345542 pir B45229 opsin, green-sensitive (clone GFgr-2)	62	2e-08
gi 4885301 ref NP 005282.1 G protein-coupled receptor 17 [_62	2e-08 5
gi 55742652 ref NP 999323.1 5-HT1D receptor [Sus scrofa] >	62	2e-08 G
gi 38016150 ref NP_937842.1 G protein-coupled receptor 103	_62	2e-08 G
gi 38016146 ref NP 937835.1 G protein-coupled receptor 103	62	2e-08 G
gi 18859537 ref NP_571661.1 vertebrate ancient long opsin gi 21307817 gb AAL25619.1 orphan G protein-coupled recepto	<u>62</u>	2e-08
gi 51860765 gb AAU11506.1 melanopsin [Phodopus sungorus]	62	2e-08
gi 14041800 dbj BAB55447.1 G protein-coupled receptor [Rat	62	2e-08 🙃
gi 47206009 emb CAF91280.1 unnamed protein product [Tetrao gi 47205825 emb CAF95884.1 unnamed protein product [Tetrao	<u>62</u>	2e-08 2e-08
gi 9823 emb CAA30644.1 rhodopsin [Octopus dofleini] >gi 12	62	2e-08
gi 25025009 ref XP 204521.1 PREDICTED: similar to trace am	62	3e-08 G
gi 21594966 gb AAH31653.1 GPR17 protein [Homo sapiens]	62	3e-08 🕒
gi 33622376 gb AAO38857.1 melanopsin [Rutilus rutilus]	62	3e-08
gi 28212246 ref NP 783180.1 trace amine receptor 14 [Rattu	62	
gi 50749504 ref XP 421666.1 PREDICTED: similar to serotoni gi 1154643 emb CAA64210.1 seretonin receptor 1D Cavia por	<u>62</u> 62	3e-08
gi 47215888 emb CAG12280.1 unnamed protein product [Tetrao	62	3e-08
gi 39591067 emb CAE58847.1 Hypothetical protein CBG02068 [62	3e-08 4e-08 G
gi 22477850 gb AAH36773.1 Opsin 3 (encephalopsin, panopsin	61	
gi 45445826 gb AAN11677.2 CG13702-PB [Drosophila melanogas	61	4e-08 G
gi 31203627 ref XP 310762.1 ENSANGP00000015565 [Anopheles gi 55620557 ref XP 526207.1 PREDICTED: similar to G protei	61 61	4e-08
gi 7657071 ref NP_055137.1 opsin 3 (encephalopsin, panopsi	61	4e-08 😘
gi 32483397 ref NP_000788.2 dopamine receptor D4 [Homo sap	61	4e-08 😉
gi 17223726 gb AAL02125.1 allatostatin C/drostatin C recep	61	4e-08
gi 291946 gb AAB59386.1 dopamine receptor D4 [Homo sapiens	<u>61</u>	4e-08 5 4e-08
gi 4325156 gb AAD17289.1 dopamine receptor D4-2 [synthetic gi 18077928 gb AAL58637.1 Dopamine D4 receptor [Homo sapiens]	61	4e-08
gi 21928798 dbj BAC05985.1 seven transmembrane helix recep	61	4e-08
gi 54641635 gb EAL30385.1 GA20386-PA [Drosophila pseudoobs	61	4e-08 4e-08
gi 47230682 emb CAF99875.1 unnamed protein product [Tetrao	01	46-00

9 of 130 $11/29/2004\ 12:32\ P\overline{N}$

Gi 47229610 emb CAG06806.1 unnamed protein product [Tetrao gi 47223619 emb CAF99228.1 unnamed protein product [Tetrao gi 47206414 emb CAF91545.1 unnamed protein product [Tetrao gi 7296517 gb AAF51802.1 CG7485-PA [Drosophila melanogaste gi 4758474 ref NP 004239.1 G protein-coupled receptor 10 [61 61 61 61	4e-08 4e-08 4e-08 5e-08
gi 27714145 ref XP 232847.1 similar to G protein-coupled r gi 29570497 gb AAO91736.1 Dopamine receptor protein 1, iso gi 8272568 gb AAF74260.1 VA opsin [Cyprinus carpio] gi 4028153 gb AAC96117.1 putative neurotransmitter recepto gi 47210163 emb CAF95187.1 unnamed protein product [Tetrao gi 47210162 emb CAF95186.1 unnamed protein product [Tetrao gi 85086 pir JH0170 octopamine receptor type I - fruit fly gi 103504 pir S12004 tyramine receptor - fruit fly (Drosop	61 61 61 61 61 61 61	5e-08 5e-08 5e-08 5e-08 5e-08 5e-08 5e-08 5e-08
gi 27685687 ref XP 220097.1 similar to G protein-coupled r gi 55664453 emb CAH73066.1 G protein-coupled receptor 10 [gi 38045882 gb AAR08905.1 nociceptin-like receptor [Rana p	60 60	7e-08 7e-08 7e-08
gi 22831755 gb AAF46059.2 CG3171-PA [Drosophila melanogast gi 55627624 ref XP 527542.1 PREDICTED: opioid receptor, mu	<u>60</u>	7e-08 G 7e-08
gi 31542912 ref NP 109658.2 G protein-coupled receptor PSPgi 992582 dbj BAA07741.1 G protein-coupled seven-transmembgi 47220968 emb CAF98197.1 unnamed protein product [Tetrao	60 60	7e-08 7e-08 7e-08
gi 11993048 gb AAG42573.1 G protein-coupled receptor PSP24 gi 639573 gb AAB30835.1 alpha 1c-adrenoceptor, alpha 1c-AR	<u>60</u>	7e-08 🗗 7e-08
gi 1122223 dbj BAA06806.1 alpha 1B adrenergic receptor [Ra gi 7707679 dbj BAA95353.1 trehalose receptor 1 [Drosophila	60 60	7e-08 ¹⁵ 7e-08

Alignments

```
Select all
                              Deselect all
 Get selected sequences
G protein-coupled receptor 78 [Homo sapiens]
gi|47480897|gb|AAH69813.1|
                          G protein-coupled receptor 78 [Homo sapiens]
 gi|47479613|gb|AAH69343.1|
 gi|34784705|gb|AAH57778.1|
                          G protein-coupled receptor 78 [Homo sapiens]
 qi|46397876|sp|Q96P69|GP78 HUMAN 🕒 Probable G protein-coupled receptor GPR78 (UNQ5925/PRO19818
 gi|21928620|dbj|BAC05898.1|
                           seven transmembrane helix receptor [Homo sapiens]
         Length = 363
 Score = 720 bits (1858), Expect = 0.0
 Identities = 363/363 (100%), Positives = 363/363 (100%)
          MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM 60
Query: 1
          MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM
Sbjct: 1
          MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM 60
Query: 61 PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP 120
          PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP
Sbjct: 61 PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP 120
Query: 121 RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG 180
          RYAGLLLGCAWGOSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG
Sbjct: 121 RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG 180
Query: 181 FVLPLAVLCLTSLOVHRVARRHCORMDTVTMKALALLADLHPSVRQRCLIQQKRRRHRAT 240
          {\tt FVLPLAVLCLTSLQVHRVARRHCQRMDTVTMKALALLADLHPSVRQRCLIQQKRRRHRAT}
Sbjct: 181 FVLPLAVLCLTSLQVHRVARRHCQRMDTVTMKALALLADLHPSVRQRCLIQQKRRRHRAT 240
Query: 241 RKIGIAIATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRP 300
          RKIGIAIATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRP
Sbjct: 241 RKIGIAIATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRP 300
Query: 301 FRQVLAGMVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQ 360
          FRQVLAGMVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQ
```

```
"Sbjct: 301 FRQVLAGMVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQ 360
   Query: 361 QTH 363
              QTH
   Sbjct: 361 QTH 363
    Length = 363
    Score = 714 \text{ bits } (1843), \text{ Expect = } 0.0
    Identities = 360/363 (99%), Positives = 361/363 (99%)
              MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM 60
              {\tt MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM}
              MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM 60
   Sbjct: 1
   Query: 61 PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP 120
              PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP
   Sbjct: 61 PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP 120
   Query: 121 RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG 180
              RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG
   Sbjct: 121 RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG 180
   Query: 181 FVLPLAVLCLTSLQVHRVARRHCQRMDTVTMKALALLADLHPSVRQRCLIQQKRRRHRAT 240
              FVLPLAVLCLTSLQVHRVARRHCQRMDTVTMKALA+LADLHPSVR CLIQQKRRRHRAT
   Sbjct: 181 FVLPLAVLCLTSLQVHRVARRHCQRMDTVTMKALAVLADLHPSVRHGCLIQQKRRRHRAT 240
   Query: 241 RKIGIAIATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRP 300
              RKIGIAIATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRP
   Sbjct: 241 RKIGIAIATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRP 300
   Query: 301 FRQVLAGMVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQ 360
              FROVLAGMVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQ
   Sbjct: 301 FRQVLAGMVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQ 360
   Query: 361 QTH 363
              OTH
   Sbjct: 361 QTH 363
    Sqi | 55622242 | ref | XP | 526521.1 | PREDICTED: similar to G protein-coupled receptor 78 [Pan
              troglodytes]
             Length = 508
    Score = 435 \text{ bits (1118)}, Expect = e-120
    Identities = 224/245 (91%), Positives = 227/245 (92%)
   Query: 1
              MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM 60
              MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM
   Sbjct: 1
              MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM 60
   Query: 61 PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP 120
              PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP
   Sbjct: 61 PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP 120
   Query: 121 RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG 180
              RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG
   Sbjct: 121 RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG 180
   Query: 181 FVLPLAVLCLTSLQVHRVARRHCQRMDTVTMKALALLADLHPSVRQRCLIQQKRRRHRAT 240
              F LPLAVLCLTSLQVHRVAR HCQRMDTVTMKALALLADLHP
                                                               Q + R
   Sbjct: 181 FALPLAVLCLTSLQVHRVARSHCQRMDTVTMKALALLADLHPRYWPSACRQAQARDLGAP 240
   Query: 241 RKIGI 245
                +G+
```

11 of 130

Sbjct: 241 WAVGL 245

```
Score = 283 bits (725), Expect = 4e-75
 Identities = 148/176 (84%), Positives = 151/176 (85%), Gaps = 7/176 (3%)
Query: 189 CLTSLQVHRVARRHCQRMDTVTMKALALLADLHP-SVRQRCLIQQKRRRHRATRKIGIAI 247
                          A+ L L P SVRQRCLIQQKRRRHRATRKIGIAI
          CL SLO
                      C
Sbjct: 339 CLPSLQPLGSGPGFCPH-----PAIILTTVLCPHSVRQRCLIQQKRRRHRATRKIGIAI 392
Query: 248 ATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRPFRQVLAG 307
          ATFLICFAPYVMTRLAELVPF+T+NAQWGILSKCLTYSKA ADPFTYSLLRRPFRQVLAG
Sbjct: 393 ATFLICFAPYVMTRLAELVPFITLNAQWGILSKCLTYSKAAADPFTYSLLRRPFRQVLAG 452
Query: 308 MVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQQTH 363
          MVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQQTH
Sbjct: 453 MVHRLLKRTPRPASTHDSSLDVAGMVHQLLKRTPRPASTHNGSVDTENDSCLQQTH 508
Si|50747354|ref|XP_426354.1| GPREDICTED: similar to G protein-coupled receptor 26 [Gallus
         Length = 416
 Score = 347 bits (890), Expect = 3e-94
 Identities = 170/299 (56%), Positives = 222/299 (74%)
         LLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDMPFTLLG 66
          LLA LLV+VL V+LLSN LVLLC YS E+R + +GV LVNLS +LLL L+MPFTLLG
Sbjct: 7
          LLALLLVLVLVVSLLSNLLVLLCFVYSTEIRKOVAGVFLVNLSFCNLLLTILNMPFTLLG 66
Query: 67 VMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRPRYAGLL 126
                   C+ +GFL+TFL SN LS+AALS D+W+AV FPL Y ++R + A +L
Sbjct: 67 ILRNQQPLGGCICKAVGFLETFLTSNTMLSMAALSIDKWIAVVFPLSYTSKMRYKDAVIL 126
Query: 127 LGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVGFVLPLA 186
          +G +W SL F +L SW+ Y+S +ASC+L L E ER RF FT H+ F+L L
Sbjct: 127 MGYSWLHSLTFPLVSLFYSWVDYNSVYASCTLHLKEETERRRFTVFTIVFHSTSFMLSLV 186
Query: 187 VLCLTSLQVHRVARRHCQRMDTVTMKALALLADLHPSVRQRCLIQQKRRRHRATRKIGIA 246
          +LC T L+V +VAR HC+R+D +TM+ L LL D+HPSV+QRCL +QKRRR RAT+KI I
Sbjct: 187 ILCFTYLKVLKVARFHCKRIDIITMQTLVLLVDIHPSVKQRCLNEQKRRRQRATKKISIF 246
Query: 247 IATFLICFAPYVMTRLAELVPFVTVNAQWGILSKCLTYSKAVADPFTYSLLRRPFRQVL 305 I +F+ICF PY++TRL EL+PFVT+N WGI+SKCLTYSKA +DPF YSLLR+ +++VL
Sbjct: 247 IGSFVICFGPYIITRLIELLPFVTINYYWGIISKCLTYSKAASDPFVYSLLRQQYKKVL 305
Length = 337
 Score = 330 bits (845), Expect = 5e-89
 Identities = 170/327 (51%), Positives = 222/327 (67%), Gaps = 4/327 (1%)
          MGPGEALLAGLLVMVLAVALLSNALVLLCCAYSAELRTRASGVLLVNLSLGHLLLAALDM 60
             +A LAGLLV + V+LLSNALVLLC +SA++R +A + +NL+ G+LL
          MNSWDAGLAGLLVGTMGVSLLSNALVLLCLLHSADIRRQAPALFTLNLTCGNLLCTVVNM 60
Sbjct: 1
Query: 61 PFTLLGVMRGRTPSAPGACQVIGFLDTFLASNAALSVAALSADQWLAVGFPLRYAGRLRP 120
          P TL GV+ R P+ C++ FLDTFLA+N+ LS+AALS D+W+AV FPL Y ++R
Sbjct: 61 PLTLAGVVAQRQPAGDRLCRLAAFLDTFLAANSMLSMAALSIDRWVAVVFPLSYRAKMRL 120
Query: 121 RYAGLLLGCAWGQSLAFSGAALGCSWLGYSSAFASCSLRLPPEPERPRFAAFTATLHAVG 180
                  W +L F AAL SWLG+ +ASC+L
                                                ER RFA FT
          R A L++
Sbjct: 121 RDAALMVAYTWLHALTFPAAALALSWLGFHQLYASCTLCSRRPDERLRFAVFTGAFHALS 180
```